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RAW SEQUENCE LISTING

DATE: 02/20/2003 TIME: 07:12:58

PATENT APPLICATION: US/09/903,190

1 <110> APPLICANT: Dumas Milne Edwards, Jean-Baptiste

Input Set : N:\Crf3\RULE60\09903190.RAW.txt Output Set: N:\CRF4\02202003\I903190.raw

Duclert, Aymeric Bouqueleret, Lydie 4 <120> TITLE OF INVENTION: Complementary DNAs 5 <130> FILE REFERENCE: GENSET.021A 6 <140> CURRENT APPLICATION NUMBER: US/09/903,190 7 <141> CURRENT FILING DATE: 2001-07-11 8 <150> PRIOR APPLICATION NUMBER: US/09/247,155A 9 <151> PRIOR FILING DATE: 1999-02-09 10 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/074,121 W--> 11 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1998-02-09 12 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/081,563 W--> 13 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-13 14 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/096,116 w--> 15 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1998-08-10 16 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/099,273 W--> 17 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1998-10-04 18 <160> NUMBER OF SEQ ID NOS: 182 19 <170> SOFTWARE: Patent.pm 21 <210> SEQ ID NO: 1 22 <211> LENGTH: 47 23 <212> TYPE: RNA 24 <213> ORGANISM: Artificial Sequence 25 <220> FEATURE: 26 <223> OTHER INFORMATION: in vitro transcription product 27 <220> FEATURE: 28 <221> NAME/KEY: modified_base 29 <222> LOCATION: 1 30 <223> OTHER INFORMATION: m7q 31 <400> SEQUENCE: 1 47 qqcauccuac ucccauccaa uuccacccua acuccuccca ucuccac 34 <210> SEO ID NO: 2 35 <211> LENGTH: 46 36 <212> TYPE: RNA 37 <213> ORGANISM: Artificial Sequence 38 <220> FEATURE: 39 <223> OTHER INFORMATION: in vitro transcription product 40 <400> SEQUENCE: 2 46 41 geauccuacu cccauccaau uccacccuaa cuccucccau cuccac 43 <210> SEQ ID NO: 3 44 <211> LENGTH: 25 45 <212> TYPE: DNA 46 <213> ORGANISM: Artificial Sequence

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		OTHER INFORMATION: oligonucleotide	
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50		atcaagaatt cgcacgagac catta	25
		SEQ ID NO: 4	
		LENGTH: 25	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
		FEATURE:	
		OTHER INFORMATION: oligonucleotide	
		SEQUENCE: 4	
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		SEQ ID NO: 5	
62	<211>	LENGTH: 25	
		TYPE: DNA	
64	<213>	ORGANISM: Artificial Sequence	
		FEATURE:	
		OTHER INFORMATION: oligonucleotide	
67	<400>	SEQUENCE: 5	
68		ccgacaagac caacgtcaag gccgc	25
		SEQ ID NO: 6	
		LENGTH: 25	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
		FEATURE:	
		OTHER INFORMATION: oligonucleotide	
76	<400>	SEQUENCE: 6	
77		tcaccagcag gcagtggctt aggag	25
		SEQ ID NO: 7	
		LENGTH: 25	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
		FEATURE:	
		OTHER INFORMATION: oligonucleotide	
		SEQUENCE: 7	
86		agtgattcct gctactttgg atggc	25
		SEQ ID NO: 8	
		LENGTH: 25	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
		FEATURE:	
		OTHER INFORMATION: oligonucleotide	
		SEQUENCE: 8	
95		gcttggtctt gttctggagt ttaga .	25
		SEQ ID NO: 9	
		LENGTH: 25	
		TYPE: DNA	
		ORGANISM: Artificial Sequence	
TOT	<220>	FEATURE:	

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102 <223> OTHER INFORMATION: oligonucleotide
     103 <400> SEQUENCE: 9
            tecagaate gagacaagee aattt
                                                                                  25
     106 <210> SEQ IF
                         10
     108
              IYPE:
     105
                        rtificial Sequence
            ORG
     11
           .0> FEA
          23> OTh NFOP
                                  jonuc
        <400> SEQ LNCF
         agggag<sup>,</sup>
                            .gcg., agtcc
                                                                                 25
            טי מדיט
      _5
                         11
      .10
                      25
      .7 <4.
                     ;NA
      18 <213> UNUANISM
     119 <220> FEATUR →
     120 < NFORMATION: oligonucleo
     1.
                   CE: 11
     1.
            u gaaagg aaaagactca tri
                                                                                 25
     12 * <210> SEQ ID N
     125 <
     126
     12' 13> 0 . NISM: Artificial Sequence
     128 <220> FEATURE:
     129 <223> OTHER INFORMATION: o gonucleo
     130 <4° SEQUENCE: 12
     131
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                                                                                 25
     13.
                  ID NO: 13
     134 . TH: 25
     135 <212> TYPE: DNA
     136 <213> ORGANISM: Artificial Sequence
     137 <220 - FEATURE:
     138 <223 · OTTER INFORMATION: oligonucleotide
     139 <400> Si
                  ™CE: 13
     140
                   t cgcacgagac catta
         at.
                                                                                 25
     142 <210> SEQ
                     : 14
     143 <211> LENGTH: 67
     144 <212> TYPE: DNA
     145 <213> ORGANISM: Artificial Sequence
     146 <220> FEATURE:
     147 <223> OTHER INFORMATION: oligonucleotide
     148 <220> FEATURE:
     149 <221> NAME/KEY: misc_feature
     150 <222> LOCATION: 67
     151 <223> OTHER INFORMATION: n=a, g, c or t
    152 <400> SEQUENCE: 14
    153
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                                                                                 60
W--> 154
              tttttvn
                                                                                 67
    156 <210> SEQ ID NO: 15
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157 <211> LENGTH: 9
158 <212> TYPE: D
159 <213> ORGANISh: Artificial Sequence
160 <220> FEATURE:
                                   cleotide
161 <223> OTHER INFORMATION: O.
162 <400> SEQUENCE: 15
                                                                                 29
       ccagcagagt cacgagagag autacau ,
     _0> SEQ ID NO: 16
165
 . .211> LF' · 25
<212> !
                 4
1.8 25 (
               : Artificial Sequence
16°
        ]
              JRE:
17. <223. OTHER INFO. ...ION: oligonucleotide
171 <400> SEOUENCE: 16
                                                                                 25
172
              agagag act:
                 YO: 17
1. .
175 <211
176 <212> TYPE: DNA
177 <213> ORGANISM: Juno Sapiens
178 <220> FEATURE:
179 <221> NAME/KEY: misc_featur_
180 <222> LOCATION: complement(261..376)
181 <223> OTHER INFORMATION: blastn
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183 <221> NAME/KEY: misc_feature
184 <222> LOCATION: complement(380..486)
185 <223> OTHER INFORMATION: blastn
186 <220> FE7 TURE:
187 <221>
          /KEY: misc_feature
            ATION: complement(110..145)
188 <222
189 <223
          THER INFORMATION: blastn
190 <220> FEATURE:
191 <221> NAME/KEY: misc_feature
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193 <223> OTHER INFORMATION: blastn
194 <220> FEATURE:
195 <221> NAME/KEY: sig_peptide
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197 <223> OTHER INFORMATION: Von Heijne matrix
198 <220> FEATURE:
199 <221> NAME/KEY: misc_feature
200 <222
            CATION: 290
201 <223
           THER INFORMATION: n=a, g, c or t
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                                                                                 60
203
          gagagaaaga actgactgar acgtttgag atg aag aaa gtt ctc ctc ctg tc
                                                                                113
204
                                          Met Lys Lys Val Leu Leu Leu 1le
205
206
          aca gcc atc ttg gca gtg gct gtw ggt ttc cca gtc tct caa gac cag
                                                                                161
207
```

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	208 209		Thr Ala Ile Leu Ala Val Ala Val Gly Phe Pro Val Ser Gln Asp Gln -5 1 5	
	210		gaa cga gaa aaa aga agt atc agt gac agc gat gaa tta gct tca ggr 209	1
	211		Glu Arg Glu Lys Arg Ser Ile Ser Asp Ser Asp Glu Leu Ala Ser Gly	
	212		10 15 20	
	213		wtt ttt gtg ttc cct tac cca tat cca ttt cgc cca ctt cca cca att 257	
M>	214		Xaa Phe Val Phe Pro Tyr Pro Tyr Pro Phe Arg Pro Leu Pro Pro Ile	
	215		25 30 35	
	216		cca ttt cca aga ttt cca tgg ttt aga cgt aan ttt cct att cca ata 305	
	217		Pro Phe Pro Arg Phe Pro Trp Phe Arg Arg Xaa Phe Pro Ile Pro Ile	
	218		40 45 50 55	
	219		cct gaa tct gcc cct aca act ccc ctt cct agc gaa aag taaacaaraa 354	
	220		Pro Glu Ser Ala Pro Thr Thr Pro Leu Pro Ser Glu Lys	
	221		60 65	
	222		ggaaaagtca crataaacct ggtcacctga aattgaaatt gagccacttc cttgaaraat 414	
	223		caaaatteet gttaataaaa raaaaacaaa tgtaattgaa atageacaca gcatteteta 474	
	224		gtcaatatct ttagtgatct tctttaataa acatgaaagc aaaaaaaaaa	
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			LENGTH: 17	
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	229	<213>	ORGANISM: Homo Sapiens	
			FEATURE:	
			NAME/KEY: SIGNAL	
			LOCATION: 117	
		<223>	OTHER INFORMATION: Von Heijne matrix	
	234		score 8.2	
	235		seq LLLITAILAVAVG/FP	
		<400>	SEQUENCE: 18	
	237		Met Lys Lys Val Leu Leu Ile Thr Ala Ile Leu Ala Val Ala Val	
	238		1 5 10 15	
	239		Gly	
			SEQ ID NO: 19	
			LENGTH: 822	
			TYPE: DNA	
			ORGANISM: Homo Sapiens	
			FEATURE:	
			NAME/KEY: misc_feature	
			LOCATION: 260464	
			OTHER INFORMATION: blastn	
			FEATURE:	
			NAME/KEY: misc_feature	
			LOCATION: 118184 OTHER INFORMATION: blastn	
			FEATURE:	
			NAME/KEY: misc_feature	
			LOCATION: 56113	
			OTHER INFORMATION: blastn	
			FEATURE:	
			NAME/KEY: misc_feature	
	200	~~~_	MANUALLE MISO_IEUCUIE	

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 02/20/2003 PATENT APPLICATION: US/09/903,190 TIME: 07:12:59

Input Set : N:\Crf3\RULE60\09903190.RAW.txt
Output Set: N:\CRF4\02202003\1903190.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:14; N Pos. 67
Seq#:17; N Pos. 290
Seg#:17; Xaa Pos. 24,50
Seq#:19; N Pos. 115
Seq#:19; Xaa Pos. 31
Seq#:21; Xaa Pos. 3
Seq#:23; N Pos. 101
Seq#:23; Xaa Pos. -2,26,27,35
Seq#:24; Xaa Pos. 14
Seq#:25; Xaa Pos. 69,106,118,134,141
Seq#:27; Xaa Pos. 156,157,170,171
Seq#:34; N Pos. 335,376
Seq#:53; N Pos. 1
Seq#:130; Xaa Pos. -5,42,58
Seq#:131; Xaa Pos. 49,74,75,88,98,104,112,114,120
Seq#:133; Xaa Pos. 48
Seq#:134; Xaa Pos. 27,28,43,44,49,50,52,53
Seq#:135; Xaa Pos. -65,31,34
Seg#:136; Xaa Pos. 48,52,55,59,65
Seq#:138; Xaa Pos. 81,84,87,131,135,143,156
Seq#:139; Xaa Pos. 32,91,98
Seq#:141; Xaa Pos. 32,100
Seq#:142; Xaa Pos. 31,84,88,90,94,100,110,113,117,122,152,153,173,193,194
Seq#:143; Xaa Pos. 37,43
Seq#:144; Xaa Pos. 56,70,107,110,113,178,181,183,195,200,202,204,230,231
Seq#:144; Xaa Pos. 244
Seq#:146; Xaa Pos. 74
Seq#:147; Xaa Pos. 4
Seq#:148; N Pos. 745
Seq#:148; Xaa Pos. -26,-25,-24
Seq#:149; N Pos. 482
Seg#:149; Xaa Pos. 40,41,43,60,70,76,82,86,105,107
Seq#:150; Xaa Pos. 59,69,78,85,87
Seq#:151; Xaa Pos. 70,139,141,154,156,165
Seq#:154; Xaa Pos. 24,25,29,87
Seq#:155; Xaa Pos. -5,42,58
Seq#:156; Xaa Pos. 49,74,75,88,98,104,112,114,120
Seq#:158; Xaa Pos. 48
Seq#:159; Xaa Pos. 27,28,43,44,49,50,52,53
Seq#:160; Xaa Pos. -65,31,34
Seg#:161; Xaa Pos. 48,52,55,59,65
Seg#:163; Xaa Pos. 81,84,87,131,135,143,156
Seq#:164; Xaa Pos. 32,91,98
Seq#:166; Xaa Pos. 32,100
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Input Set : N:\Crf3\RULE60\09903190.RAW.txt
Output Set: N:\CRF4\02202003\I903190.raw

Seq#:167; Xaa Pos. 31,84,88,90,94,100,110,113,117,122,152,153,173,193,194

Seq#:168; Xaa Pos. 37,43

 $\bar{\text{Seq\#:169; Xaa Pos.}} \ 56,70,107,110,113,178,181,183,195,200,202,204,230,231$

Seq#:169; Xaa Pos. 244
Seq#:171; Xaa Pos. 74
Seq#:172; Xaa Pos. 4

Seq#:173; Xaa Pos. -26,-25,-24

VERIFICATION SUMMARY PATENT APPLICATION: US/09/903,190 DATE: 02/20/2003 TIME: 07:12:59

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L:11 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:13 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:15 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:17 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:60
L:214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:257
M:341 Repeated in SeqNo=17
L:295 \text{ M}:341 \text{ W}: \text{ (46) "n" or "Xaa" used, for SEQ ID$$\#:19 after pos.:60}
M:341 Repeated in SeqNo=19
L:361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:325
L:412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:60
M:341 Repeated in SeqNo=23
L:447 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:478 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:345
M:341 Repeated in SeqNo=25
L:552~M:341~W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:580
M:341 Repeated in SeqNo=27
L:909 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:300
M:341 Repeated in SeqNo=34
L:1842 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 after pos.:0
L:4994 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:130 after pos.:98
M:341 Repeated in SeqNo=130
L:5052 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131 after pos.:298
M:341 Repeated in SeqNo=131
L:5166 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:133 after pos.:404
L:5218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:134 after pos.:407
M:341 Repeated in SeqNo=134
L:5253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:135 after pos.:49
M:341 Repeated in SeqNo=135
L:5315 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:136 after pos.:246
M:341 Repeated in SeqNo=136
L:5411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138 after pos.:394
M:341 Repeated in SeqNo=138
L:5471 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139 after pos.:254
M:341 Repeated in SeqNo=139
L:5588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:141 after pos.:288
M:341 Repeated in SeqNo=141
L:5637 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:142 after pos.:198
M:341 Repeated in SeqNo=142
L:5718 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:143 after pos.:341
L:5771 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:144 after pos.:291
M:341 Repeated in SeqNo=144
L:5906 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146 after pos.:352
L:5983 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:147 after pos.:316
L:6022 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:148 after pos.:162
M:341 Repeated in SeqNo=148
L:6083 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:149 after pos.:371
M:341 Repeated in SeqNo=149
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VERIFICATION SUMMARY

DATE: 02/20/2003

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TIME: 07:12:59

Input Set : N:\Crf3\RULE60\09903190.RAW.txt Output Set: N:\CRF4\02202003\I903190.raw

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M:341 Repeated in SeqNo=150

 $L:6200 \ M:341 \ W:$ (46) "n" or "Xaa" used, for SEQ ID#:151 after pos.:349

M:341 Repeated in SeqNo=151

L:6420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154 after pos.:296

M:341 Repeated in SeqNo=154